

Figure 1

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M 1

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E H P L F G C L R S P H A T A Q G L H P F S Q S S 26

CTG GCC CTC CAT GGA AGA TCT GAC CAC ATG TCC TAC CCC GAA CTC TCC ACA TCT TCC TCG TCT TGC ATA ATC GCG 349
L A L H G R S D H H S Y P E L S T S S S S C I I A 51

GGA TAC CCC AAT GAG GAG GGC ATG TTT GCC AGC CAG CAT CAC AGC GCG CAC CAC CAC CAC CAC CAC CAC CAC CAT 424
G Y P N E E G H F A S Q H H R G H H H H H H H H H H 76

CAC CAC CAC CAG CAG CAG CAG CAG GCT CTG CAA AGC AAC TGG CAC CTC CCG CAG ATG TCC TCC CCG CCA AGC 499
H H H Q Q Q Q Q H Q A L Q S N W H L P Q M S S P P S 101

GCG GCC CGG CAC AGC CTT TGC CTG CAG CCT GAT TCC GGA GGG CCC CCG GAG CTG GGG AGC AGC CCT CCG GTC CTG 574
A A R H S L C L Q P D S G G P P E L G S S P P V L 126

TGC TCC AAC TCT TCT AGC CTG GGC TCC AGC ACC CCG ACC GGA GCC GCG TGC GCA CCA AGG GAT TAT GGC CGT CAA 649
C S N S S S L G S S T P T G A A C A P R D Y G R Q 151

GCG CTG TCA CCC GCA GAA GTG GAG AAG AGA AGT GGC AGC AAA AGA AAA AGC GAC AGT TCA GAT TCC CAG GAA GGA 724
A L S P A E V E K R S G S K R K S D S S D S Q E G 176

AAT TAC AAG TCA GAA GTG AAC AGC AAA CCT AGG AAG GAA AGA ACA GCT TTC ACC AAA GAG CAA ATC AGA GAA CTT 799
N Y K S E V N S K P R K Z R T A F T K Z Q I R E L 201

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GAA AGA CAG GTG AAA GTG TGG TTC CAG AAC AGG AGA ATG AAG TGG AAG CCG GTC AAG GGG GGA CAA CAA GGA GCT 949
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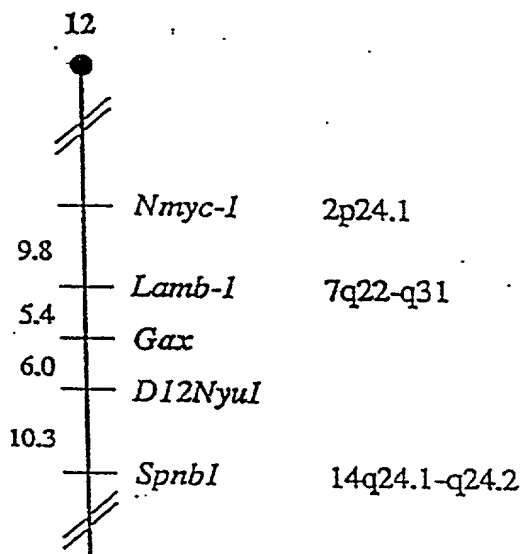
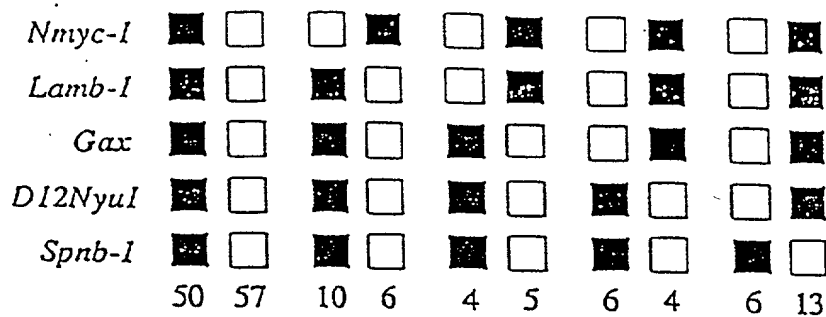
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Figure 2

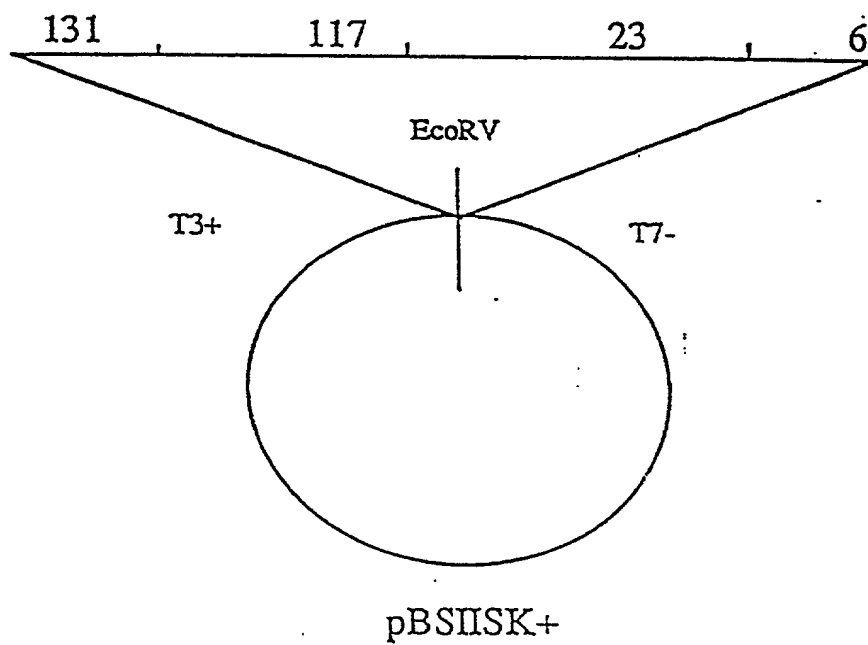


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30																			
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56	G	L	H	P	F	S	Q	S	L	A	L	H	Q	R	S	D	H	N	S
69																			
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96	T	S	S	S	S	C	I	I	A	G	Y	P	N	E	E	D	M	F	A
109																			
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269																			
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Figure 3

Figure 4



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Figure 5a

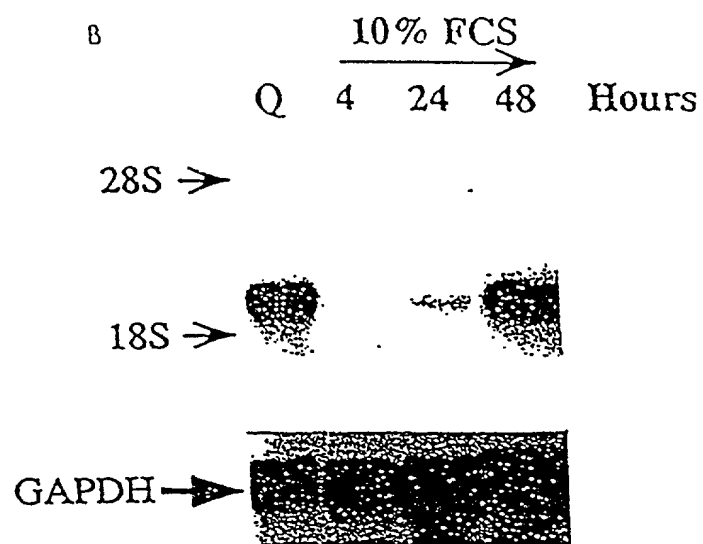
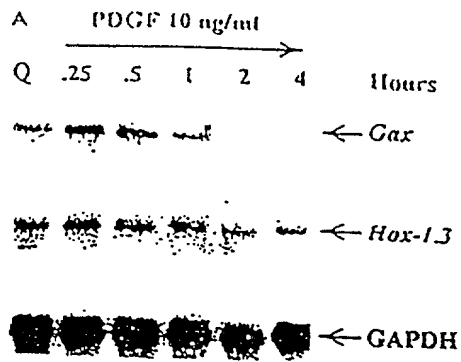
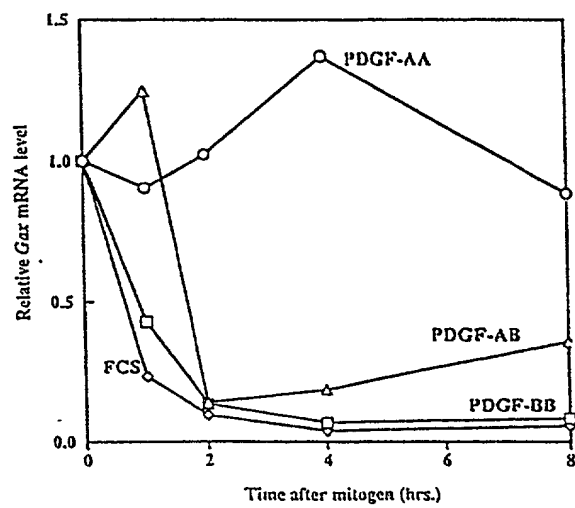


Figure 5B



0394067.082704

Figure 6



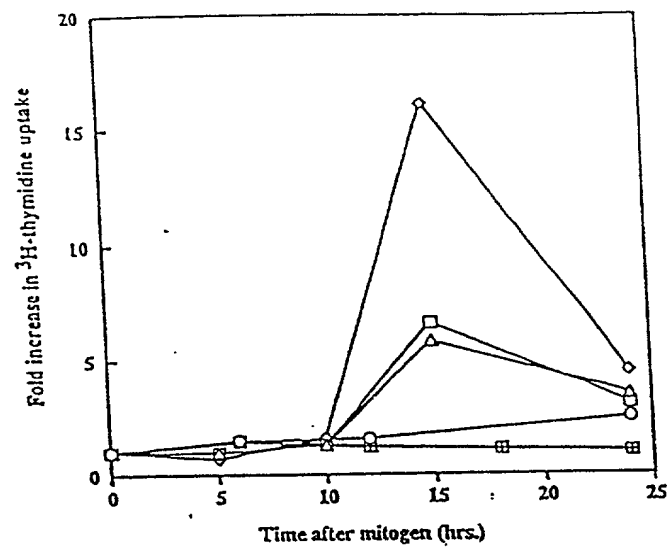


Figure 7

Figure 8

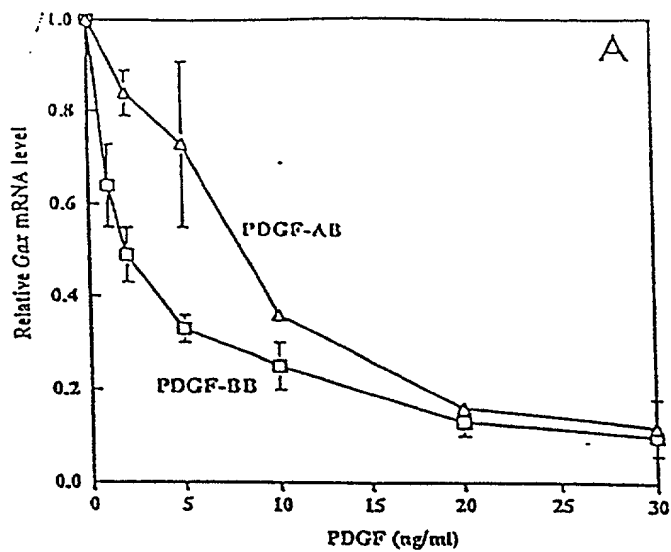


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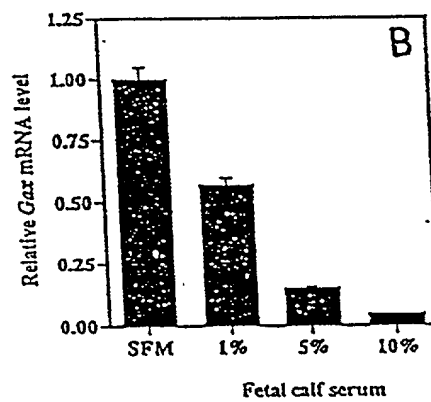
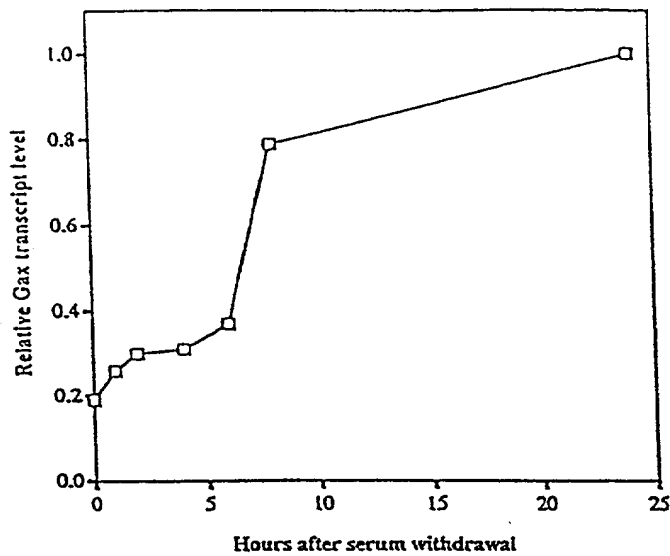


Figure 10



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Figure 11

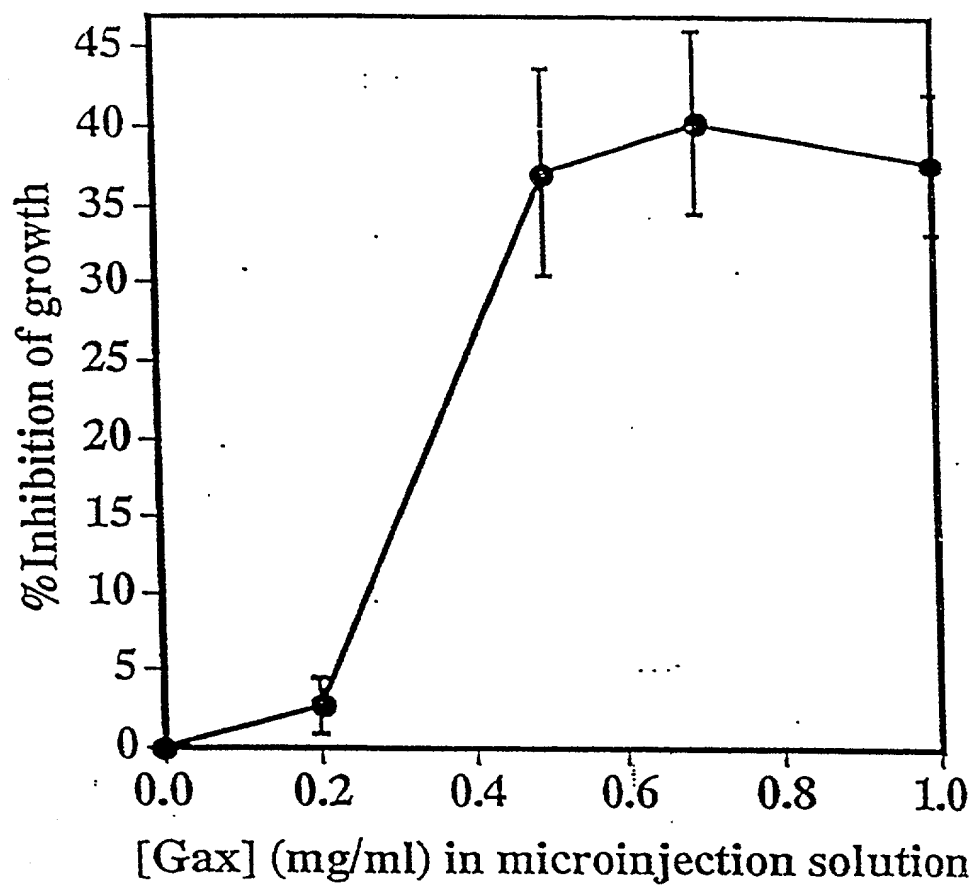


Figure 10

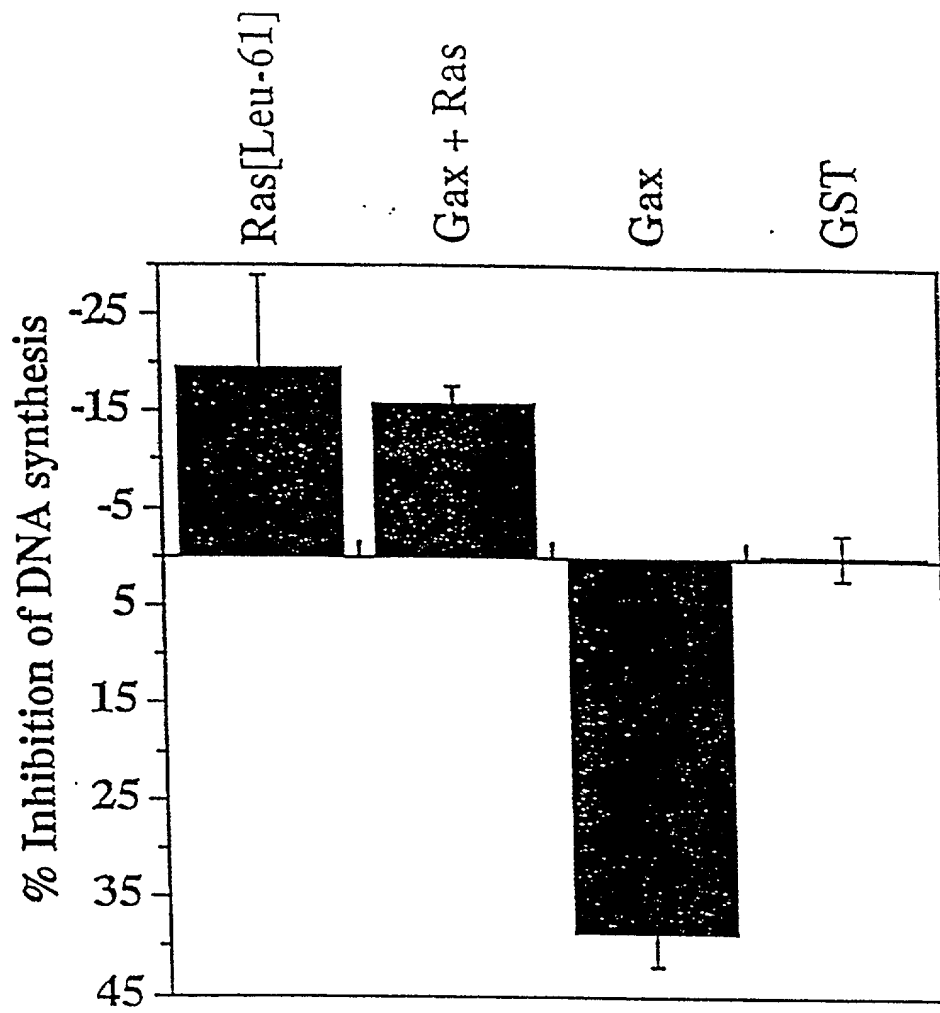
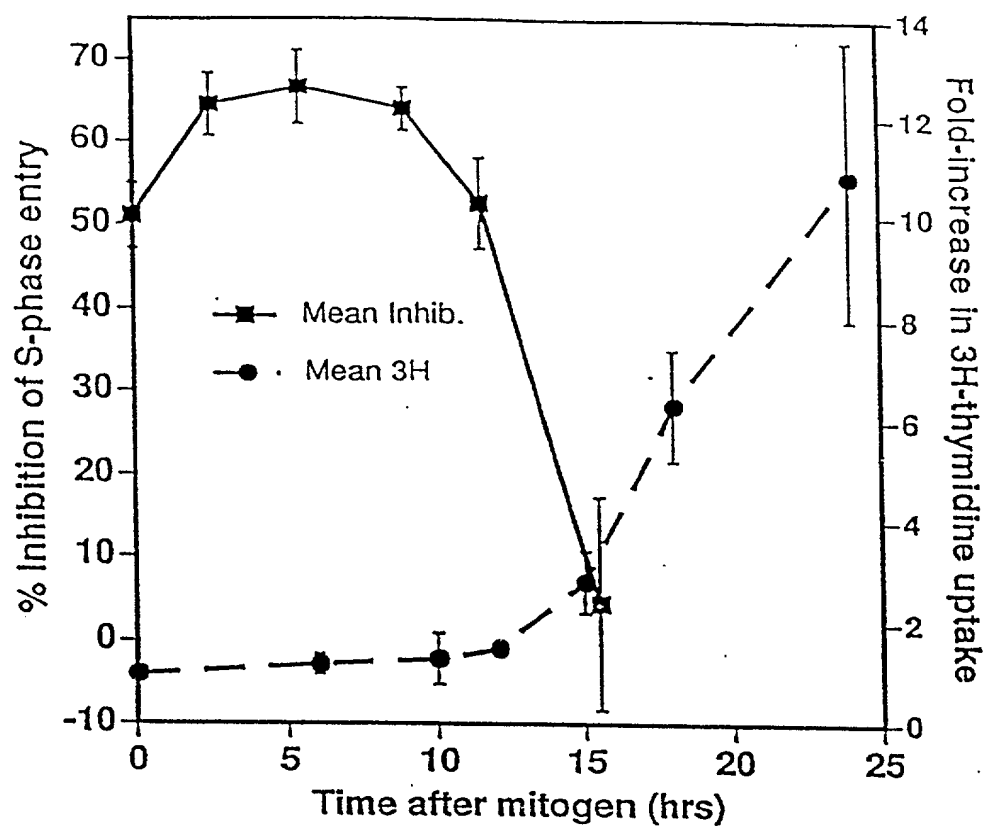
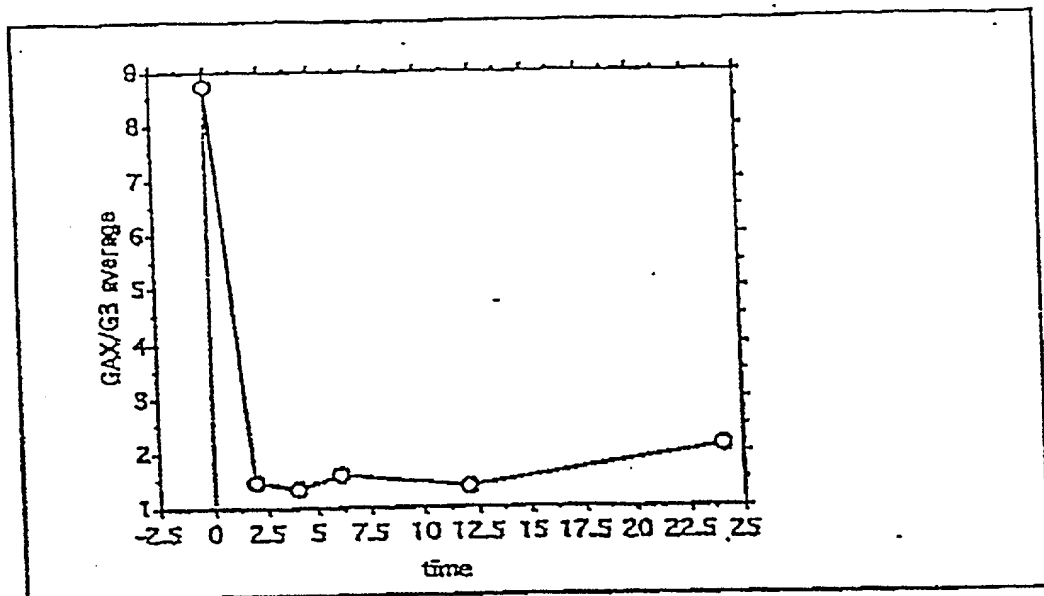


Figure 13B



05940673.022701
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Figure 14



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